

MLLNVLRICI	IVCLVNDGAG	KHSEGRERTK	TYSLNSRGYF	40
RKERGARRSK	ILLVNTKGLD	EPHIGHGDFG	LVAELFDSTR	80
THTNRKEPDM	NKVKLFSTVA	HGNKSARRKA	YNGSRRNIFS	120
RRSFDKRNTE	VTEKPGAKMF	WNNFLVKMNG	APQNTSHGSK	160
AQEIMKEACK	TLPFTQNIVH	ENCDRMVIQN	NLCFGKCISL	200
HVPNQODRRN	TCSHCLPSKF	TLNHLTLNCT	GSKNVVKVVM	240
MVEECTCEAH	KSNFHQTAQF	NMDTSTTLHH		270

Figure 1. Deduced amino acid sequence of *Xenopus cerberus* protein. SEQ ID NO:1.

Figure 2. Nucleotide sequence of the full-length cerberus DNA derived from the Xenopus organizer. The sense strand is on top (in the 5' to 3' direction) and the antisense strand on the bottom line (on the opposite direction). SEQ ID NO:2.

GAATTC	CCCAG	CAAGTC	GCTC	AGAAAC	ACTG	CAGGGT	CTAG	ATATCA	TACA	ATGTTA	CTAA	60
CTTAAG	GGTC	GTTCAG	CGAG	TCTTTG	TGAC	GTCCAG	ATC	TATAGT	TATGT	TACAAT	GATT	
ATGTA	CTCAG	GATCTG	TATT	ATCGT	CTGCC	TTGTGA	ATGA	TGGAG	CAGGA	AAAC	ACTCAG	120
TACAT	GAGTC	CTAGAC	ATAA	TAGCAG	ACGG	AACACT	TACT	ACCTCG	TCT	TTGTG	AGTC	
AAGGAC	GAGA	AAGGAC	AAAA	ACATAT	TCAC	TTAAC	AGCAG	AGGTTA	CTTC	AGAAA	AGAAA	180
TTCTG	CTCT	TTCTG	TTTT	TGTATA	AGTG	AATTGT	CGTC	TCCAAT	GAAG	TCTTT	TCTTT	
GAGGAG	CACG	TAGGAG	CAAG	ATTCTG	CTGG	TGAATA	CTAA	AGGTCT	TGAT	GAACCC	CACA	240
CTCCT	CGTG	ATCCT	CGTT	TAAGAC	GACC	ACTTAT	GATT	TCCAGA	AACTA	CTTGGG	TGT	
TTGGG	CATGG	TGATTT	TCGC	TTAGTA	AGCTG	AAC	TATTTGA	TTCCAC	CAGA	ACACAT	ACAA	300
AACCCG	TACC	ACTAAA	AGCG	AATCAT	CGAC	TTGATA	AACT	AAGGTG	GCT	TGTGT	ATGTT	
ACAGAA	AAGA	GCCAG	ACATG	AACAA	AGTCA	AGCTTT	TCTC	AACAGT	TGCC	CATGGA	AAACA	360
TGCTTT	TTCT	CGGTCT	GTAC	TTGTTT	TCAGT	TCGAAA	AGAG	TTGTCA	ACGG	GTACCT	TTGT	
AAAGT	GCAAG	AAGAAA	AGCT	TACAAT	TGTT	CTAGA	AGGAA	TATTTT	TCCT	CGCCGT	TCTT	420
TTTCA	CGTT	TTCTTT	TCGA	ATGTTA	CCAA	GATCTT	CCCT	ATAAAA	AGGA	GCGGCA	AGAA	
TTGATA	AAAG	AAATAC	AGAG	GTTACT	GAAA	AGCCTG	GTGC	CAAGAT	GTTC	TGGAAC	AATT	480
AACTAT	TTTC	TTTATG	TCTC	CAATGA	CTTT	TCGGAC	CACG	GTTCTA	CAAG	ACCTGT	TAA	
TTTTG	GTAA	AATGA	ATGGA	GCCCCA	CAGA	ATACA	AGCCA	TGGCAG	TAAA	GCACAG	GAAA	540
AAAAC	CAATT	TTACTT	ACCT	CGGGGT	GTCT	TATGTT	CGGT	ACCGT	CATT	CGTGT	CCTTT	
TAATGA	AAGA	AGCTTG	CAAA	ACCTGT	TTTT	TTACTC	AGAA	TATTGT	TACAT	GAAA	ACTGTG	600
ATTACT	TTCT	TCGAAC	GTTT	TGGAAC	AAAA	AGTGAG	TCTT	ATAAC	ATGTA	CTTTT	GACAC	
ACAGG	ATGGT	GATAC	AGAAC	AATCTG	TGCT	TTGGTA	AAATG	CATCT	CTCTC	CATGTT	CCAA	660
TGTCCT	TACCA	CTATGT	CTTG	TTAGAC	CACGA	AACCAT	TTAC	GTAGAG	AGAG	GTACA	AGGTT	
ATCAG	CAAGA	TCGAC	GAAAT	ACTTGT	TCCC	ATTGCT	TGCC	GTCCAA	ATTT	ACCTG	AACC	720
TAGTC	GTTCT	AGCTG	CTTTA	TGAACA	AGGG	TAACGA	ACGG	CAGGTT	TAAA	TGGG	ACTTGG	
ACCTG	ACGCT	GAATTG	TACT	GGATCT	TAAGA	ATGTAG	TAAA	GGTTGT	CATG	ATGGT	AGAGG	780
TGGAC	TGCGA	CTTAAC	ATGA	CCTAG	ATTCT	TACAT	CATT	CCAAC	AGTAC	TACCAT	CTCC	
AATGC	ACGTG	TGAAG	CTCAT	AAGAG	CAACT	TCCAC	CAAAC	TGCAC	AGTTT	AACAT	GGATA	840
TTACG	TGCAC	ACTTC	GAGTA	TTCTC	GTTGA	AGGTG	GTTT	ACGTG	TCAA	TTGT	TACCTAT	
CATCT	ACTAC	CCTGC	ACCAT	TAAAG	GA	CTG	CTGAC	GGTAT	GTCAT	ACCTT	ACGG	900
GTAGAT	GTATG	GGACG	TGGTA	ATTC	CTGAC	GGTAT	GTCAT	ACCTT	ACGG	GAAA	ACAACC	
AATAT	TTTGT	ACATA	CTATG	CATCT	AAAGC	ATTAT	GTTGC	CTTCT	ATTT	ATATA	ACCAC	960
TTATA	AAACAA	TGTAT	GATAC	GTAGA	TTTCG	TAATA	CAACG	GAAGAT	AAAG	TATAT	TGGTG	
ATGGA	ATAAG	GATTG	TATGA	ATTATA	ATTA	ACAAAT	TGGCA	TTTTG	TGTAA	CATG	CAAGAT	1020
TACCT	TATTC	CTAAC	ATACT	TAATAT	TAAAT	TGTTT	TACCGT	AAAAC	ACATT	GTACG	TCTA	

MSRTRKVDL LLLAIPGLAL LLLPNAYCAS CEPVRIPMCK SMPWNMTKMP NHLHHSTQAN	60
AILAIEQFEG LLTTECSQDL LFFLCAMYAP ICTIDFQHEP IKPCKSV CER ARAGCEPILI	120
KYRHTWPESL ACEELPVYDR GVCISPEAIV TVEQGTDSMP DFSMDSNNGN CGSGREHCKC	180
KPMKATQKTY LKNNYNYVIR AKVKEVKVVC HDATAIVEVK EILKSSLVNI PKDTVTLYTN	240
SGCLCPQLVA NEEYIIMGYE DKERTRLLLV EGSLAEKWRD RLAKKVVRWD QKLRRPRKSK	300
DPVAPIPNKN SNSRQARS	

Figure 3. Deduced amino acid sequence of Xenopus frazzled protein. SEQ ID NO:3.

Figure 4. Nucleotide sequence of the full-length frazzled cDNA derived from the *Xenopus* organizer. The sense strand of the DNA on top (5' to 3' direction) and the antisense strand on the bottom line (opposite direction). SEQ ID NO:4.

GAATTCCCTT TCACACAGGA CTCCTGGCAG AGGTGAATGG TTAGCCCTAT GGATTGTT	60
CTTAAGGGAA AGTGTGTCCT GAGGACCGTC TCCACTTACC AATCGGGATA CCTAAACCA	
TGTTGATTTT GACACATGAT TGATTGCTTT CAGATAGGAT TGAAGGACTT GGATTTTAT	120
ACAACTAAA CTGTGTACTA ACTAACGAAA GTCTATCCTA ACTTCCTGAA CCTAAAAATA	
CTAATTCCTGC ACTTTTAAAT TATCTGAGTA ATTGTTCAAT TTGTATTGGA TGGGACTAAA	180
GATTAAGACG TGAAAATTTA ATAGACTCAT TAACAAGTAA AACATAACCT ACCCTGATTT	
GATAAACTTA ACTCCTTGCT TTTGACTTGC CCATAAACTA TAAGGTGGGG TGAGTTGTAG	240
CTATTTGAAT TGAGGAACGA AAAGTGAACG GGTATTTGAT ATTCCACCCC ACTCAACATC	
TTGCTTTTAC ATGTGCCAG ATTTTCCCTG TATTCCCTGT ATTCCCTCTA AAGTAAGCCT	300
AACGAAAATG TACACGGGTC TAAAAGGGAC ATAAGGGACA TAAGGGAGAT TTCATTGCGA	
ACACATACAG GTTGGGCAGA ATAACAATGT CTCGAACAAG GAAAGTGGAC TCATTACTGC	360
TGTGTATGTC CAACCCGTCT TATTGTTACA GAGCTTGTTT CTTTCACCTG AGTAATGACG	
TACTGGCCAT ACCTGGACTG GCGCTTCTCT TATTACCCAA TGCTTACTGT GCTTCGTGTG	420
ATGACCGGTA TGGACCTGAC CGCGAAGAGA ATAATGGGTT ACGAATGACA CGAAGCACAC	
AGCCTGTGCG GATCCCCATG TGCAAACTA TGCCATGGAA CATGACCAAG ATGCCCAACC	480
TCGGACACGC CTAGGGGTAC ACGTTTAGAT ACGGTACCTT GTACTGGTTC TACGGGTTGG	
ATCTCCACCA CAGCACTCAA GCCAATGCCA TCCTGGCAAT TGAACAGTTT GAAGGTTTGC	540
TAGAGGTGGT GTCGTGAGTT CGGTTACGGT AGGACCGTTA ACTTGTCAAA CTTCCAAACG	
TGACCACTGA ATGTAGCCAG GACCTTTTGT TCTTTCTGTG TGCCATGTAT GCCCCATTT	600
ACTGGTGACT TACATCGGTC CTGGAACA AGAAGACAC ACGGTACATA CGGGGGTAAA	
GTACCATCGA TTTCCAGCAT GAACCAATTA AGCCTTGCAA GTCCGTGTGC GAAAGGGCCA	660
CATGGTAGCT AAAGGTCGTA CTTGGTTAAT TCGGAACGTT CAGGCACACG CTTTCCGGT	
GGGCCGGCTG TGAGCCCAT CTCATAAAGT ACCGGCACAC TTGGCCAGAG AGCCTGGCAT	720
CCCGGCCGAC ACTCGGTAA GAGTATTTCA TGGCCGTGTG AACCAGTCTC TCGGACCGTA	
GTGAAGAGCT GCGCGTATAT GACAGAGGAG TCTGCATCTC CCCAGAGGCT ATCGTCACAG	780
CACCTTCTCGA CGGGCATATA CTGTCTCCTC AGACGTAGAG GGGTCTCCGA TAGCAGTGTC	
TGGAACAAGG AACAGATTCA ATGCCAGACT TCTCCATGGA TTCAAACAAT GGAAATTGCG	840
ACCTTGTTCC TTGTCTAAGT TACGGTCTGA AGAGGTACCT AAGTTTGTTA CCTTTAACGC	
GAAGCGGCAG GGAGCACTGT AAATGCAAGC CCATGAAGGC AACCCAAAAG ACGTATCTCA	900
CTTCGCCGTC CCTCGTGACA TTTACGTTTC GGTACTTCCG TTGGGTTTTT TGCAATAGAGT	
AGAATAATTA CAATTATGTA ATCAGAGCAA AAGTGAAAGA GGTGAAAGTG AAATGCCACG	960
TCTTATTAAT GTTAATACAT TAGTCTCGTT TTCACCTTCT CCACTTTCAC TTTACGGTGC	
ACGCAACAGC AATTGTGGAA GTAAAGGAGA TTCTCAAGTC TTCCCTAGTG AACATTCCTA	1020
TGCGTTGTGC TTAACACCTT CATTTCTCT AAGAGTTCAG AAGGGATCAC TTGTAAGGAT	

AGACACACAGT	GACACTGTAC	ACCAACTCAG	GCTGCTTTGTG	CCCCACGCTT	GTTGCCAATG	1080
TTCTGTGTCA	CTGTGACATG	TGGTTGAGTC	CGACGAACAC	GGGGGTCGAA	CAACGGTTAC	
AGGAATACAT	AATTATGGGC	TATGAAGACA	AAGAGCGTAC	CAGGCTTCTA	CTAGTGGAAG	1140
TCCTTATGTA	TTAATACCCG	ATACTTCTGT	TTCTCGCATG	GTCCGAAGAT	GATCACCTTC	
GATCCTTGGC	CGAAAAATGG	AGAGATCGTC	TTGCTAAGAA	AGTCAAGCGC	TGGGATCAAA	1200
CTAGGAACCG	GCTTTTTTACC	TCTCTAGCAG	AACGATTCTT	TCAGTTCGCG	ACCCTAGTTT	
AGCTTCGACG	TCCCAGGAAA	AGCAAAGACC	CCGTGGCTCC	AATTCCCAAC	AAAAACAGCA	1260
TCGAAGCTGC	AGGGTCCTTT	TCGTTTCTGG	GGCACCAGAG	TTAAGGGTTG	TTTTTGTCGT	
ATTCCAGACA	AGCGCGTAGT	TAGACTAACG	GAAAGGTGTA	TGGAAACTCT	ATGGACTTTG	1320
TAAGGTCTGT	TCGCGCATCA	ATCTGATTGC	CTTTCACAT	ACCTTTGAGA	TACCTGAAAC	
AAACTAAGAT	TTGCATTGTT	GGAAGAGCAA	AAAAGAAATT	GCACTACAGC	ACGTTATATT	1380
TTTGATTCTA	AACGTAACAA	CCTTCTCGTT	TTTTCTTTAA	CGTGATGTCG	TGCAATATAA	
CTATTGTTTA	CTACAAGAAG	CTGGTTTAGT	TGATTGTAGT	TCTCCTTTCC	TTCTTTTTTT	1440
GATAACAAAT	GATGTTCTTC	GACCAAATCA	ACTAACATCA	AGAGGAAAGG	AAGAAAAAAA	
TTATAACTAT	ATTTGCACGT	GTTCCCAGGC	AATTGTTTTA	TTCAACTTCC	AGTGACAGAG	1500
AATATTGATA	TAAACGTGCA	CAAGGGTCCG	TTAACAAAAT	AAGTTGAAGG	TCACTGTCTC	
CAGTGACTGA	ATGTCTCAGC	CTAAGAAGC	TCAATTCATT	TCTGATCAAC	TAATGGTGAC	1560
GTCACTGACT	TACAGAGTCG	GATTTCTTCG	AGTTAAGTAA	AGACTAGTTG	ATTACCACTG	
AAGTGTTTGA	TACTTGGGGA	AAGTGAACTA	ATTGCAATGG	TAAATCAGAG	AAAAGTTGAC	1620
TTCACAAAC	ATGAACCCCT	TTCACTTGAT	TACGTTTACC	ATTTAGTCTC	TTTTCAACTG	
CAATGTTGCT	TTTCTGTAG	ATGAACAAGT	GAGAGATCAC	ATTTAAATGA	TGATCACTTT	1680
GTTACAACGA	AAAGGACATC	TACTTGTTC	CTCTCTAGTG	TAAATTTACT	ACTAGTGAAA	
CCATTTAATA	CTTTCAGCAG	TTTTAGTTAG	ATGACATGTA	GGATGCACCT	AAATCTAAAT	1740
GGTAAATTAT	GAAAGTCGTC	AAAATCAATC	TACTGTACAT	CCTACGTGGA	TTTAGATTTA	
ATTTTATCAT	AAATGAAGAG	CTGGTTTAGA	CTGTATGGTC	ACTGTTGGGA	AGGTAAATGC	1800
TAAATAGTA	TTTACTTCTC	GACCAAATCT	GACATACCAG	TGACAACCC	TCCATTACG	
CTACTTTGTC	AATTCTGTTT	TAAAAATTGC	CTAAATAAAT	ATTAAGTCCT	AAATAAAAAA	1860
GATGAAACAG	TTAAGACAAA	ATTTTAAACG	GATTTATTTA	TAATTCAGGA	TTTATTTTTT	
AAAAAAAAAA	AAAAA					
TTTTTTTTTT	TTTTT					

Fig. 4. (Continuation page 2, SEQ ID NO:4).

MLLLFRAIPM LLLGLMVLQT DCEIAQYYID EEEPPGTVIA VLSQHSIFNT TDIPATNFRL 60
 MKQFNNSLIG VRESQGQLSI MERIDREQIC RQSLHCNLAL DVVSFSKGHF KLLNVKVEVR 120
 DINDHSPHFP SEIMHVEVSE SSSVGTRIPL EIAIDEDVGS NSIQNFQISN NSHFSIDVLT 180
 RADGVKYADL VLMRELDREI QPTYIMELLA MDGGVPSLSG TAVVNIRVLD FNDNSPVFER 240
 STIAVDLVED APLGYLLEL HATDDDEGVN GEIVYGFSTL ASQEVRLFK INSRTGSVTL 300
 EGQVDFETKQ TYEFEVQAQD LGPNPLTATC KVTVHILDVN DNTPAITITP LTTVNAGVAY 360
 IPETATKENF IALISTDRA SGSNGQVRCT LYGHEHFKLQ QAYEDSYMIV TTSTLDRENI 420
 AAYSLTVVAE DLGFPSLTKK KYITVKVSDE NDNAPVFSKP QYEASILENN APGSYITTVI 480
 ARSDSDQNG KVNRYRLVDAK VMGQSLTTFV SLDADSGVLR AVRSLDYEKL KQLDFEIEAA 540
 DNGIPQLSTR VQLNLRIVDQ NDNCPVITNP LLNNGSGEVL LPISAPQNYL VFQLKAEDSD 600
 EGHNSQLFYT ILRDPSRLFA INKESGEVFL KKQLNSDHSE DLSIVVAVYD LGRPSLSTNA 660
 TVKFILTDSF PSNVEVVILQ PSAEEQHQID MSIIFIAVLA GGCALLLLAI FFVACTCKKK 720
 AGEFKQVPEQ HGTCNEERLL STPSPQSVSS SLSQSESCQL SINTESENCV VSSNQEQHQQ 780
 TGIKHSISVP SYHTSGWHLN NCAMSISGHS HMGHISTKVQ WAKEIVTSMT VTLILVENQK 840
 RRALSSQCRH KPVLTNTQMNQ QGSDMPITIS ATESTRVQKM GTAHCNMKRA IDCLTL

Figure 5. Deduced amino acid sequence of the *Xenopus* PAPC (paraxial protocadherin) protein. It encodes a member of the cadherin family of transmembrane proteins that has dorsalizing activity when constructs are injected into *Xenopus* embryos. SEQ ID NO:5.

Figure 6. Nucleotide sequence of the full-length PAPC cDNA derived from the *Xenopus* organizer. The sense strand of the DNA is shown in the top line (in the 5' to 3' direction), and the bottom line shows the antisense strand (opposite orientation). SEQ ID NO:6.

GAATTC	CCAG	AGATGA	ACTC	CTGAGATTG	TTTAAATGA	CTGCAGGTCT	GGAAGGATTC	60		
CTTAAG	GGGTC	TCTACTT	GAG	GA	ACTCTAAC	AAAATTTACT	GACGTCCAGA	CCTTCCTAAG		
ACATTG	CCAC	ACTGTTT	CTA	GGCATG	AAAA	A	ACTGCAAGT	TTCAACTTTG	TTTTTGGTGC	120
TGTAAC	GGTG	TGACAA	AGAT	CCGTACT	TTTT	TTGACGTTCA	AAGTTGAAAC	AAAA	ACCACG	
AACTTG	GATT	CTTCAAG	ATG	CTGCTT	CTCT	TCAGAGCCAT	TCCAATGCTG	CTGTTGGGAC	180	
TTGAA	ACTAA	GAAGTT	CTAC	GACGAAG	GAGA	AGTCTCGGTA	AGGTTACGAC	GACA	ACCCTG	
TGATGG	TTTT	ACAAAC	GAGAC	TGTGAA	AATTG	CCCAGTACTA	CATAGATGAA	GAAGA	ACCCC	240
ACTAC	CAAAA	TGTTT	GTCTG	ACACTT	TAAAC	GGGTCATGAT	GATCTACTT	CTTCT	TGGGG	
CTGGC	ACTGT	AATTGC	AGTG	TTGTC	CACAAC	ACTCCATATT	TAACACTACA	GATATA	CCCTG	300
GACCGT	GACA	TTAACG	TCAC	AACAGT	GTTG	TGAGGTATAA	ATTGTGATGT	CTATAT	GAGAC	
CAACCA	ATTT	CCGTCT	AAATG	AAGCA	ATTTA	ATAATTCCCT	TATCGGAGTC	CGTGAG	AGTG	360
GTTGGT	TAAA	GGCAG	ATTAC	TTCGT	TAAAT	TATTAAGGGA	ATAGCCTCAG	GCACT	CTCAC	
ATGGG	CAGCT	GAGCAT	CATG	GAGAGG	ATTG	ACCGGGAGCA	AATCTGCAGG	CAGTCC	CTTC	420
TACCCG	TCGA	CTCGT	AGTAC	CTCTC	CTAAC	TGGCCCTCGT	TTAGACGTCC	GTCAGG	GGAAG	
ACTGCA	ACCT	GGCTTT	GAT	GTGGT	CAGCT	TTTCCAAAGG	ACACTTCAAG	CTTCTG	AACG	480
TGACGT	TGGA	CCGAA	ACCTA	CACCA	GTCA	AAAGGTTTCC	TGTGAAGTTC	GAAGAC	TGTC	
TGAAAG	TGGA	GGTGAG	AGAC	ATTAAT	GACC	ATAGCCCTCA	CTTTCCAGT	GAAATA	ATGC	540
ACTTTC	CACCT	CCACT	CTCTG	TAATT	ACTGG	TATCGGGAGT	GAAAGGGTCA	CTTTAT	TACG	
ATGTG	GAGGT	GTCTG	AAAGT	TCTCT	GTGG	GCACCAGGAT	TCCTTTAGAA	ATTGCA	ATAG	600
TACAC	CTCCA	CAGACT	TTTCA	AGGAG	ACACC	CGTGGTCCTA	AGGAAATCTT	TAACGT	TATC	
ATGAAG	ATGT	TGGGT	CCAAC	TCCAT	CCAGA	ACTTTCAGAT	CTCAAATAAT	AGCCAC	TCTCA	660
TACTTC	TACA	ACCCAG	GTTG	AGGTAG	GTCT	TGAAAGTCTA	GAGTTTATTA	TCGGT	GAAAT	
GCATT	GATGT	GCTAAC	CCAGA	GCAGAT	TGGG	TGAAATATGC	AGATTTAGTC	TTAATG	AGAG	720
CGTAA	CTACA	CGATT	GCTCT	CGTCT	ACCCC	ACTTTATACG	TCTAAATCAG	AATTACT	CTC	
AACTGG	ACAG	GGAAAT	CCAG	CCAAC	ATACA	TAATGGAGCT	ACTAGCAATG	GATGGG	GGTG	780
TTGAC	CTGTC	CCTTT	AGGTC	GGTTG	TATGT	ATTACCTCGA	TGATCGTTAC	CTACCC	CCAC	
TACCAT	CACT	ATCTG	GTTACT	GCAGT	GGTTA	ACATCCGAGT	CCTGGACTTT	AATGATA	ACA	840
ATGGT	AGTGA	TAGAC	CATGA	CGTCA	CCAAT	TGTAGGCTCA	GGACCTGAAA	TTACTAT	TGT	
GCCAG	TGTT	TGAGAG	AAGC	ACCATT	GTCTG	TGGACCTAGT	AGAGGATGCT	CCTCTG	GGAT	900
CGGGT	CACAA	ACTCT	CTTTCG	TGGTA	ACGAC	ACCTGGATCA	TCTCCTACGA	GGAGAC	CCTA	
ACCTTT	TGTT	GGAGT	TACAT	GCTACT	GACG	ATGATGAAGG	AGTGAATGGA	GAAATT	GTTT	960
TGGA	AAACAA	CCTCA	ATGTA	CGATG	ACTGC	TACTACTTCC	TCACTTACCT	CTTTAA	CAAA	
ATGG	ATTCAG	CACTTT	TGGCA	TCTCA	AGAGG	TACGTCAGCT	ATTTAAAATT	AACTCC	AGAA	1020
TACCTA	AGTC	GTGAA	ACCGT	AGAGT	TCTCC	ATGCAGTCGA	TAAATTTTAA	TTGAGG	TCTT	

[illegible]

Fig. 6. (Continuation page 3, SEQ ID NO:6).

ATTAAATCCA CAGACCTACA GTCAAATATT TGAGGGCCCC TGAAACAGCA CATCAGTCAG 3360
 TAATTTAGGT GTCTGGATGT CAGTTTATAA ACTCCCGGGG ACTTTGTCGT GTAGTCAGTC

 GACCTAAAGT GGCCTTTTTA CTTTTAGCAG CTCCTGGGTC TGCCCTCTGT GTTAATCAGC 3420
 CTGGATTTC CCGGAAAAAT GAAAATCGTC GAGGACCCAG ACGGGAGACA CAATTAGTCG

 CCCTGGTCAA GTCCTGAGTA GGATCATGGC GTTTTATAT GCATCTCACC TACTTTGGAC 3480
 GGGACCAGTT CAGGACTCAT CCTAGTACCG CAAAAATATA CGTAGAGTGG ATGAAACCTG

 GTGATTACA CATAATAGGA AACGCTTGGT TTCAGTGAAG TCTGTGTTGT ATATATTCTG 3540
 CACTAAATGT GTATTATCCT TTGCGAACCA AAGTCACTTC AGACACAACA TATATAAGAC

 TTATATACAC GCATTTTGTG TTTGTGTATA TATTCAAGT CCATTAGAT ATGTGTATAT 3600
 AATATATGTG CGTAAACAC AAACACATAT ATAAAGTTCA GGTAAGTCTA TACACATATA

 AGTGCAGACC TTGTAAATTA AATATTCTGA TACTTTTCC TCAATAAATA TTTAAAT
 TCACGTCTGG AACATTTAAT TTATAAGACT ATGAAAAGG AGTTATTAT AAATTTA

Fig. 6. (Continuation page 4, SEQ ID NO:6).

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MVCCGPGRML LGWAGLLVLA ALCLLQVPGA QAAACEPVRI PLCKSLPWNM TKMPNHLHHS 60
TQANAILAME QFEGLLGTHC SPDLLFFLCA MYAPICTIDF QHEPIKPCKS VCERARQGCE 120
PILIKYRHSW PESLACDELP VYDRGVCISP EAIVTADGAD FPMSSTGHC RGASSERCKC 180
KPV RATQKTY FRNNYNYVIR AKVKEVKMKC HDVTAVVEVK EILKASLVNI PRDTVNLYTT 240
SGCLCPPLTV NEEYVIMGYE DEERSRLLLV EGSIAEKWKD RLGKKVKRWD MKLRHLGLGK 300
TDASDSTQNQ KSGRNSNPRP ARS.

Figure 7. Deduced amino acid sequence of mouse FRZB-1 protein. SEQ ID NO:7.

FRZB-1

Variable	Mean	SD	Min	Max
Age	38.5	12.5	20	65
Gender	0.5	0.5	0	1
Marital status	0.5	0.5	0	1
Education	12.5	1.5	9	16
Income	1500	500	500	3000
Health status	0.5	0.5	0	1
Smoking status	0.5	0.5	0	1
Alcohol consumption	0.5	0.5	0	1
Exercise frequency	0.5	0.5	0	1
Stress level	0.5	0.5	0	1
Sleep quality	0.5	0.5	0	1
Appetite	0.5	0.5	0	1
Weight change	0.5	0.5	0	1
Blood pressure	120	10	90	150
Cholesterol level	200	30	150	250
Glucose level	100	10	70	130
Hemoglobin A1c	5.5	0.5	4.5	6.5
Insulin sensitivity	0.5	0.5	0	1
Diabetes risk score	0.5	0.5	0	1
Overall health score	0.5	0.5	0	1

AAGCCTGGGA TTCGGACCCCT	CCATGGTCTG GGTACCAGAC	CTGCGGCCCG GACGCCGGGC	GGACGGATGC CCTGCCTACG	TGCTAGGATG ACGATCCTAC	GGCCGGGTTG CCGCCCCAAC	60
CTAGTCCTGG GATCAGGACC	CTGCTCTCTG GACGAGAGAC	CCTGCTCCAG GGACGAGGTC	GTGCCCCGGAG CACGGGCCTC	CTCAGGCTGC GAGTCCGACG	AGCCTGTGAG TCGGACACTC	120
CCTGTCCGCA GGACAGGCGT	TCCCCGCTGTG AGGGCGACAC	CAAGTCCCTT GTTCAGGGAA	CCCTGGAACA GGGACCTTGT	TGACCAAGAT ACTGGTTCTA	GCCCAACCAC CGGGTTGGTG	180
CTGCACCACA GACGTGGTGT	GCACCCAGGC CGTGGGTCCG	TAACGCCATC ATTGCGGTAG	CTGGCCATGG GACCGGTACC	AACAGTTCGA TTGTCAAGCT	AGGGCTGCTG TCCCACGAC	240
GGCACCCTACT CCGTGGGTGA	GCAGCCCGGA CGTCGGGCCT	TCTTCTCTTC AGAAGAGAAG	TTCCTCTGTG AAGGAGACAC	CAATGTACGC GTTACATGCG	ACCCATTTGC TGGGTAAACG	300
ACCATCGACT TGGTAGCTGA	TCCAGCACGA AGGTCGTGCT	GCCCATCAAG CGGGTAGTTC	CCCTGCAAGT GGGACGTTCA	CTGTGTGTGA GACACACACT	GCGCGCCCGA CGCGCGGGCT	360
CAGGGCTGCG GTCCCACGCG	AGCCCATTTCT TCGGGTAAGA	CATCAAGTAC GTAGTTCATG	CGCCACTCGT GCGGTGAGCA	GGCCGGAAAG CCGGCCTTTC	CTTGGCCTGC GAACCGGACG	420
GACGAGCTGC CTGCTCGACG	CGGTGTACGA GCCACATGCT	CCGCGGCGTG GGCGCCGCAC	TGCATCTCTC ACGTAGAGAG	CTGAGGCCAT GACTCCGGTA	CGTCACCGCG GCAGTGGCGC	480
GACGGAGCGG CTGCCTCGCC	ATTTTCCCTAT TAAAAGGATA	GGATTCAAGT CCTAAGTTCA	ACTGGACACT TGACCTGTGA	GCAGAGGGGC CGTCTCCCCG	AAGCAGCGAA TTCGTGCTTT	540
CGTTGCAAAT GCAACGTTTA	GTAAGCCTGT CATTCGGACA	CAGAGCTACA GTCTCGATGT	CAGAAGACCT GTCTTCTGGA	ATTTCCGGAA TAAAGGCCTT	CAATTACAAC GTTAATGTTG	600
TATGTCATCC ATACAGTAGG	GGGCTAAAGT CCCGATTTCA	TAAAGAGGTA ATTTCTCCAT	AAGATGAAAT TTCTACTTTA	GTCTATGATG CAGTACTACA	GACCGCCGTT CTGGCGGCAA	660
GTGGAAGTGA CACCTTCACT	AGGAAATTCT TCCTTTAAGA	AAAGGCATCA TTTCCGTAGT	CTGGTAAACA GACCATTTGT	TTCCAAGGGA AAGGTTCCCT	CACCGTCAAT GTGGCAGTTA	720
CTTTATACCA GAAATATGGT	CCTCTGGCTG GGAGACCGAC	CCTCTGTCCT GGAGACAGGA	CCACTTACTG GGTGAATGAC	TCAATGAGGA AGTTACTCCT	ATATGTCATC TATACAGTAG	780
ATGGGCTATG TACCCGATAC	AAGACGAGGA TTCTGTCTCT	ACGTTCCAGG TGCAAGGTCC	TTACTCTTGG AATGAGAACC	TAGAAGGCTC ATCTTCCGAG	TATAGCTGAG ATATCGACTC	840
AAGTGGAAGG TTCACCTTCC	ATCGGCTTGG TAGCCGAACC	TAAGAAAGTC ATTCTTTTCA	AAGCGCTGGG TTCGCGACCC	ATATGAAACT TATACTTTGA	CCGACACCTT GGCTGTGGAA	900
GGACTGGGTA CCTGACCCAT	AAACTGATGC TTTGACTACG	TAGCGATTCC ATCGCTAAGG	ACTCAGAATC TGAGTCTTAG	AGAAGTCTGG TCTTCAGACC	CAGGAACTCT GTCCTTGAGA	960

AATCCCCGGC	CAGCACGCAG	CTAAATCCTG	AAATGTAAAA	GGCCACACCC	ACGGACTCCC	1020
TTAGGGGCCG	GTCGTGCGTC	GATTTAGGAC	TTTACATTTT	CCGGTGTGGG	TGCCTGAGGG	
TTCTAAGACT	GGCGCTGGTG	GACTAACAAA	GGAAACCGC	ACAGTTGTGC	TCGTGACCGA	1080
AAGATTCTGA	CCGCGACCAC	CTGATTGTTT	CCTTTTGGCG	TGTCAACACG	AGCACTGGCT	
TTGTTTACCG	CAGACACCGC	GTGGCTACCG	AAGTTACTTC	CGGTCCCCTT	TCTCCTGCCT	1140
AACAAATGGC	GTCTGTGGCG	CACCGATGGC	TTCAATGAAG	GCCAGGGGAA	AGAGGACGAA	
CTTAATGGCG	TGGGGTTAGA	TCCTTTAATA	TGTTATATAT	TCTGTTTCAT	CAATCACGTG	1200
GAATTACCGC	ACCCCAATCT	AGGAAATTAT	ACAATATATA	AGACAAAGTA	GTTAGTGCAC	
GGGACTGTTC	TTTTGCAACC	AGAATAGTAA	ATTAAATATG	TTGATGCTAA	GGTTTCTGTA	1260
CCCTGACAAG	AAAACGTTGG	TCTTATCATT	TAATTTATAC	AACTACGATT	CCAAAGACAT	
CTGGACTCCC	TGGGTTTAAT	TTGGTGTTCT	GTACCCTGAT	TGAGAATGCA	ATGTTTCATG	1320
GACCTGAGGG	ACCCAAATTA	AACCACAAGA	CATGGGACTA	ACTCTTACGT	TACAAAGTAC	
TAAAGAGAGA	ATCCTGGTCA	TATCTCAAGA	ACTAGATATT	GCTGTAAGAC	AGCCTCTGCT	1380
ATTTCTCTCT	TAGGACCAGT	ATAGAGTTCT	TGATCTATAA	CGACATTCTG	TCGGAGACGA	
GCTGCGCTTA	TAGTCTTGTG	TTTGTATGCC	TTTGTCCATT	TCCCTCATGC	TGTGAAAGTT	1440
CGACGCGAAT	ATCAGAACAC	AAACATACGG	AAACAGGTAA	AGGGAGTACG	ACACTTTCAA	
ATACATGTTT	ATAAAGGTAG	AACGGCATT	TGAAATCAGA	CACTGCACAA	GCAGAGTAGC	1500
TATGTACAAA	TATTTCCATC	TTGCCGTAAA	ACTTTAGTCT	GTGACGTGTT	CGTCTCATCG	
CCAACACCAG	GAAGCATTTA	TGAGGAAACG	CCACACAGCA	TGACTTATTT	TCAAGATTGG	1560
GGTTGTGGTC	CTTCGTAAAT	ACTCCTTTGC	GGTGTGTCGT	ACTGAATAAA	AGTTCTAACC	
CAGGCAGCAA	AATAAATAGT	GTTGGGAGCC	AAGAAAAGAA	TATTTTGCCT	GGTTAAGGGG	1620
GTCCGTCGTT	TTATTTATCA	CAACCCTCGG	TTCTTTTCTT	ATAAAACGGA	CCAATTCCCC	
CACACTGGAA	TCAGTAGCCC	TTGAGCCATT	AACAGCAGTG	TTCTTCTGGC	AAGTTTTTTGA	1680
GTGTGACCTT	AGTCATCGGG	AACTCGGTAA	TTGTCGTCAC	AAGAAGACCG	TTCAAAAAC	
TTTGTTTCATA	AATGTATTCA	CGAGCATTAG	AGATGAACTT	ATAACTAGAC	ATCTGTTGTT	1740
AAACAAGTAT	TTACATAAGT	GCTCGTAATC	TCTACTTGAA	TATTGATCTG	TAGACAACAA	
ATCTCTATAG	CTCTGCTTCC	TTCTAAATCA	AACCCATTGT	TGGATGCTCC	CTCTCCATTTC	1800
TAGAGATATC	GAGACGAAGG	AAGATTTAGT	TTGGGTAACA	ACCTACGAGG	GAGAGGTAAG	

ATAAATAAAAT	TTGGCTTGCT	GATATTGGCCA	GGAAAAAGAAA	GTATTAAAGT	ATGCATGCAT	1860
TATTTATTTA	AACCGAACGA	CATAACCGGT	CCTTTTCTTT	CATAATTTCA	TACGTACGTA	
GTGCACCAGG	GTGTTATTTA	ACAGAGGTAT	GTAACCTCTAT	AAAAGACTAT	AATTTACAGG	1920
CACGTGGTCC	CACAAATAAA	TGTCTCCATA	CATTGAGATA	TTTTCTGATA	TTAAATGTCC	
ACACGGAAAT	GTGCACATTT	GTTTACTTTT	TTTCTTCCTT	TTGCTTTGGG	CTTGTGATTT	1980
TGTGCCTTTA	CACGTGTAAA	CAAAATGAAAA	AAAGAAGGAA	AACGAAACCC	GAACACTAAA	
TGGTTTTTTGG	TGTGTTTATG	TCTGTATTTT	GGGGGGTGGG	TAGGTTTAAG	CCATTGCACA	2040
ACCAAAAACC	ACACAAATAC	AGACATAAAA	CCCCCACCCC	ATCCAAATTC	GGTAACGTGT	
TTCAAGTTGA	ACTAGATTAG	AGTAGACTAG	GCTCATTTGGC	CTAGACATTA	TGATTTGAAT	2100
AAGTTCAACT	TGATCTAATC	TCATCTGATC	CGAGTAACCG	GATCTGTAAT	ACTAAACTTA	
TTGTGTTGTT	TAATGCTCCA	TCAAGATGTC	TAATAAAAAGG	AATATGGTTG	TCAACAGAGA	2160
AACACAACAA	ATTACGAGGT	AGTTCTACAG	ATTATTTTCC	TTATACCAAC	AGTTGTCTCT	
CGACAACAAC	AACAAA					
GCTGTTGTTG	TTGTTT					

[illegible]

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Figure 10. Nucleotide sequence of the full-length human FRZB-1 cDNA. SEQ ID NO:10.

This sequence was assembled from public ESTs from the Genbank database

(accession numbers: H18848, R63748, W38677, W44760, H38379 and N71244).

GGCGGAGCGG GCCTTTTGGC GTCCACTGCG CGGCTGCACC CTGCCCCATC TGCCGGGATC 60
CCGCCTCGCC CGGAAAACCG CAGGTGACGC GCCGACGTGG GACGGGGTAG ACGGCCCTAG

ATGGTCTGCG GCAGCCCGGG AGGGATGCTG CTGCTGCGGG CCGGGCTGCT TGCCCTGGCT 120
TACCAGACGC CGTCGGGCCC TCCCTACGAC GACGACGCCC GGCCCGACGA ACGGGACCGA

GCTCTCTGCC TGCTCCGGGT GCCCGGGGCT CGGGCTGCAG CCTGTGAGCC CGTCCGCATC 180
CGAGAGACGG ACGAGGCCCA CGGGCCCCGA GCCCGACGTC GGACACTCGG GCAGGCGTAG

CCCCTGTGCA AGTCCCTGCC CTGGAACATG ACTAAGATGC CCAACCACCT GCACCACAGC 240
GGGGACACGT TCAGGGACGG GACCTTGATC TGATTCTACG GGTGTTGGTA CGTGGTGTCTG

ACTCAGGCCA ACGCCATCCT GGCCATCGAG CAGTTCGAAG GTCTGCTGGG CACCCACTGC 300
TGAGTCCGGT TGCGGTAGGA CCGGTAGCTC GTCAAGCTTC CAGACGACCC GTGGGTGACG

AGCCCCGATC TGCTCTTCTT CCTCTGTGCC ATGTACGCGC CCATCTGCAC CATTGACTTC 360
TCGGGGCTAG ACGAGAAGAA GGAGACACGG TACATGCGCG GGTAGACGTG GTAAGTGAAG

CAGCACGAGC CCATCAAGCC CTGTAAGTCT GTGTGCGAGC GGGCCCCGCA GGGCTGTGAG 420
GTCGTGCTCG GGTAGTTCGG GACATTCAGA CACACGCTCG CCCGGGCCGT CCCGACACTC

CCCATACTCA TCAAGTACCG CCACTCGTGG CCGGAGAACC TGGCCTGCGA GGAGCTGCCA 480
GGGTATGAGT AGTTCATGGC GGTGAGCACC GGCCTCTTGG ACCGGACGCT CCTCGACGGT

GTGTACGACA GGGGCGTGTG CATCTCTCCC GAGGCCATCG TTAGTGCGGA CGGAGCTGAT 540
CACATGCTGT CCGCGCACAC GTAGAGAGGG CTCGGGTAGC AATGACGCCT GCCTCGACTA

TTTCCTATGG ATTCTAGTAA CGGAAACTGT AGAGGGGCAA GCAGTGAACG CTGTAAATGT 600
AAAGGATACC TAAGATCATT GCCTTTGACA TCTCCCCGTT CGTCACTTGC GACATTTACA

AAGCCTATTA GAGCTACACA GAAGACCTAT TTCCGGAACA ATTACAATA TGTCATTCGG 660
TTCGGATAAT CTCGATGTGT CTTCTGGATA AAGGCCTTGT TAATGTTGAT ACAGTAAGCC

GCTAAAGTTA AAGAGATAAA GACTAAGTGC CATGATGTGA CTGCAGTAGT GGAGGTGAAG 720
CGATTTCAAT TTCTCTATTT CTGATTCACG GTACTACACT GACGTCATCA CCTCCACTTC

GAGATTCTAA AGTCCTCTCT GGTAACATT CCACGGGACA CTGTCAACCT CTATACCAGC 780
CTCTAAGATT TCAGGAGAGA CCATTTGTAA GGTGCCCTGT GACAGTTGGA GATATGGTCTG

TCTGGCTGCC TCTGCCCTCC ACTTAATGTT AATGAGGAAT ATATCATCAT GGGCTATGAA 840
AGACCGACGG AGACGGGAGG TGAATTACAA TTAATCCTTA TATAGTAGTA CCCGATACTT

GATGAGGAAC	GTTCAGATT	ACTCTGGTG	GAAGGCTCTA	TAGCTGAGAA	GTGGAAGGAT	900
CTACTCCTTG	CAAGGTCTAA	TGAGAACCAC	CTTCCGAGAT	ATCGACTCTT	CACCTTCCTA	
CGACTCGGTA	AAAAAGTTAA	GCGCTGGGAT	ATGAAGCTTC	GTCATCTTGG	ACTCAGTAAA	960
GCTGAGCCAT	TTTTTCAATT	CGCGACCCTA	TACTTCGAAG	CAGTAGAACC	TGAGTCATTT	
AGTGATTCTA	GCAATAGTGA	TTCCACTCAG	AGTCAGAAGT	CTGGCAGGAA	CTCGAACCCC	1020
TCACTAAGAT	CGTTATCACT	AAGGTGAGTC	TCAGTCTTCA	GACCGTCCTT	GAGCTTGGGG	
CGGCAAGCAC	GCAACTAAAT	CCCGAAATAC	AAAAAGTAAC	ACAGTGGACT	TCCTATTAAG	1080
GCCGTTTCGTG	CGTTGATTTA	GGGCTTTATG	TTTTTTCATTG	TGTCACCTGA	AGGATAATTC	
ACTTACTTGC	ATTGCTGGAC	TAGCAAAGGA	AAATTGCACT	ATTGCACATC	ATATTCTATT	1140
TGAATGAACG	TAACGACCTG	ATCGTTTCCT	TTTAACGTGA	TAACGTGTAG	TATAAGATAA	
GTTTACTATA	AAAATCATGT	GATAACTGAT	TATTACTTCT	GTTTCTCTTT	TGGTTTCTGC	1200
CAAATGATAT	TTTTAGTACA	CTATTGACTA	ATAATGAAGA	CAAAGAGAAA	ACCAAAGACG	
TTCTCTCTTC	TCTCAACCCC	TTTGTAAATGG	TTTGGGGGCA	GACTCTTAAG	TATATTGTGA	1260
AAGAGAGAAG	AGAGTTGGGG	AAACATTACC	AAACCCCCGT	CTGAGAATTC	ATATAACACT	
GTTTTCTATT	TCACTAATCA	TGAGAAAAAC	TGTTCTTTTG	CAATAATAAT	AAATTAAACA	1320
CAAAAGATAA	AGTGATTAGT	ACTCTTTTTG	ACAAGAAAAC	GTTATTATTA	TTTAATTTGT	
TGCTGTTACC	AGAGCCTCTT	TGCTGAGTCT	CCAGATGTTA	ATTTACTTTT	TGCACCCCCA	1380
ACGACAATGG	TCTCGGAGAA	ACGACTCAGA	GGTCTACAAT	TAAATGAAAG	ACGTGGGGTT	
TTGGGAATGC	AATATTGGAT	GAAAAGAGAG	GTTTCTGGTA	TTACACAGAA	GCTAGATATG	1440
AACCCTTACG	TTATAACCTA	CTTTTCTCTC	CAAAGACCAT	AAGTGCTTTT	CGATCTATAC	
CCTTAAAACA	TACTCTGCCG	ATCTAATTAC	AGCCTTATTT	TTGTATGCCT	TTTGGGCATT	1500
GGAATTTTGT	ATGAGACGGC	TAGATTAATG	TCGGAATAAA	AACATACGGA	AAACCCGTAA	
CTCCTCATGC	TTAGAAAAGTT	CCAAATGTTT	ATAAAAGGTAA	AATGGCAGTT	TGAAGTCAAA	1560
GAGGAGTACG	AATCTTTCAA	GGTTTACAAA	TATTTCCATT	TTACCGTCAA	ACTTCAGTTT	
TGTCACATAG	GCAAAGCAAT	CAAGCACCAG	GAAGTGTTTA	TGAGGAAACA	ACACCCAAGA	1620
ACAGTGTATC	CGTTTCGTTA	GTTCTGGTTC	CTTCACAAAT	ACTCCTTTGT	TGTGGGGTCT	
TGAATTATTT	TTGAGACTGT	CAGGAAGTAA	AATAAATAGG	AGCTTAAGAA	AGAACATTTT	1680
ACTTAATAAA	AACCTTGACA	GTCCTTCATT	TTATTTATCC	TCGAATTCCT	TCTTGTAATA	
GCCTGATTGA	GAAGCACAAAC	TGAAACCAGT	AGCCGCTGGG	GTGTTAATGG	TAGCATTCCT	1740
CGGACTAACT	CTTCGTGTTG	ACTTTGGTCA	TCGGCGACCC	CACAATTACC	ATCGTAAGAA	
CTTTTGGCAA	TACATTTGAT	TTGTTTCATGA	ATATATTAAT	CAGCATTAGA	GAAATGAATT	1800
GAAACCGTT	ATGTAAACTA	AACAAGTACT	TATATAATTA	GTCGTAATCT	CTTTACTTAA	
ATAACTAGAC	ATCTGCTGTT	ATCACCATAG	TTTTGTTTAA	TTTGCTTCCT	TTTAAATAAA	1860
TATTGATCTG	TAGACGACAA	TAGTGGTATC	AAAACAAATT	AAACGAAGGA	AAATTTATTT	
CCCATTTGGTG	AAAGTCAAAA	AAAAAAAAAA	AAA			
GGGTAACCAC	TTTCAGTTTT	TTTTTTTTTT	TTT			